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Regult
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Maximum DB
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                  1046
1044.5
678.5
677.5
631.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Published Applications AA:*

1: /cgn2_6/ptodata/1/pubpaa/US07 PUBCOMB.pep:*
2: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
3: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
4: /cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
5: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
6: /cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
7: /cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
9: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
9: /cgn2_6/ptodata/1/pubpaa/US09B_PUBCOMB.pep:*
10: /cgn2_6/ptodata/1/pubpaa/US09B_PUBCOMB.pep:*
11: /cgn2_6/ptodata/1/pubpaa/US09C_PUBCOMB.pep:*
13: /cgn2_6/ptodata/1/pubpaa/US09C_PUBCOMB.pep:*
14: /cgn2_6/ptodata/1/pubpaa/US09C_PUBCOMB.pep:*
15: /cgn2_6/ptodata/1/pubpaa/US09C_PUBCOMB.pep:*
16: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
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18: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
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length: 2000000000
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Gapop 10.0 , Gapext 0.5
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Listing first 45 summaries
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     US-09-758-017A-2
US-09-758-017A-4
4 US-10-038-010-12
4 US-10-106-698-5713
5 US-10-369-493-2360
5 US-10-369-493-23507
5 US-10-269-877-14
5 US-10-269-877-14
5 US-10-269-877-14
5 US-10-369-493-1593
5 US-10-369-493-15818
5 US-10-369-493-13816
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5 US-10-369-493-13816
5 US-10-369-493-13816
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2152.432 Million cell updates/sec
Sequence 2, Appli
Sequence 4, Appli
Sequence 12, Appl
Sequence 2713, Appl
Sequence 216, Appl
Sequence 21107, A
Sequence 23507, A
Sequence 14, Appl
Sequence 1593, A
Sequence 1593, A
Sequence 11818, A
Sequence 11886, A
Sequence 10331, A
Sequence 11885, A
Sequence 11885, A
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e 27601 e 18891	Sequence 9066, Ap Sequence 1562, Ap	e 8212,	Sequence 322, App Sequence 4961, Ap	e 18312	Sequence 13321, A	Sequence 188911,	0	8693,	19129	4570,	e 823,	7327,	22663	1860,	14539	23353	17469	17831	9434,	15044	e 10737	e 17754	e 9179,	e 16617	

ALIGNMENTS

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APPLICANT: Lanes, Olav
APPLICANT: Lanes, Olav
APPLICANT: Lanes, Nils Peder
APPLICANT: Gidlasen, Nils Peder
APPLICANT: Gidlasen, Nils Peder
APPLICANT: Gidlasen, Nils Peder
TITLE OF INVENTION: Cod uracil-DNA glycosylase, gene coding therefore,
TITLE OF INVENTION: recombinant DNA containing said gene or operative parts
TITLE OF INVENTION: thereof, a method for preparing said protein and the
TITLE OF INVENTION: use of said protein or preparing said protein and the
TITLE OF INVENTION use of said protein or said operative pa
TITLE OF INVENTION NUMBER: US/09/758,017A
CURRENT APPLICATION NUMBER: 2000 5428
PRIOR APPLICATION NUMBER: 2000 5428
PRIOR APPLICATION NUMBER: 2000 10-32
PRIOR APPLICATION NUMBER: 2000 10-3
PRIOR APPLICATION NUMBER: 2000 5428
PRIOR FILING DATE: 2000-01-12
NUMBER OF SEQ ID NOS: 19
SEQ ID NO 2
LENCTH- 301
                                                                                                                                                                                                                                                                                                    ; LENGTH: 301
; TYPE: PRT
; ORGANISM: Gadus morhua
US-09-758-017A-2
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                                                                                                                                                                                          Query Match 99.0%;
Best Local Similarity 99.3%;
Matches 299; Conservative
                               19
RMAKVKKAALDKIRAKATPAGFGETWRRELAAEFEKPYFKQLMSFVADERSRHTVYPPAD
                                                                                                               MLFKLGLCQRCISSNRXLPGLLIPQTLCFSKLMKITPKKLRSSNVEQKTSSPQLSVEQLE 60
                                                                                      MLFKLGLCQRCISSNRVLPGLLIPQTLCFSKLMKITPKKLRSSNVEQKTSSPQLSVEQLE 60
                                                                                                                                                                                          Score 1591; DB 9;
Pred. No. 4.5e-161;
0; Mismatches 2;
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APPLICANT: Lanes, Olav
APPLICANT: Willasen, Nils Peder
APPLICANT: Willasen, Nils Peder
APPLICANT: Giddal, Per Henrik
APPLICANT: Gjellesvik, Dag Rune
TITLE OF INVENTION: Cod uracil-DNA glycosylase, gene coding therefore,
TITLE OF INVENTION: recombinant DNA containing said gene or operative part
TITLE OF INVENTION: thereof, a method for preparing said protein and the
TITLE OF INVENTION: use of said protein or said operative pa
TITLE OF INVENTION use of said protein or said operative pa
TITLE OF INVENTION UMBER: US/09/758,017A
CURRENT APPLICATION NUMBER: 2001-01-10
PRIOR APPLICATION NUMBER: 2000-5428
PRIOR APPLICATION NUMBER: 2000 0163
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US-09-758-017A-4
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SEQ ID NO 4
LENGTH: 301
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 2000-01-12
NUMBER OF SEQ ID NOS: 19
 273
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                                                                                                                                                                                                                                                                                                                                                      Similarity 95.;
56; Conservative
                RGFLGCKHFSKANGLLKLSGTEPINWRAL 301
                                                                                    SHKDRGWETFTDAVIKWLSVNREGVVFLLWGSYAHKKGATIDRKRHHVLQAVHPSPLSAH
                                                                                                                                                                                                                                                                              VQTTPKKLRSSNVEQKTSSPQLSVEQLERMAKNKKAALDKTRAKATPAGFGETWRRELAA
                                                                    SHKDRGWETSTDAVIKWLSVNREGVVFLFWGSYAHKKGATIDRKRHHVLQALHPSPLSAH
                                                                                                                                      LCFSVQKPVPPPPSLVNIYKELCTDIDGFKHPGHGDLSGWANKGCCCLTRCLTVRAHQAN
                                                                                                                                                            LCFSVQKPVPPPPSLVNIYKELCTDIDGFKHPGHGDLSGWAKQGVLLLNAVLTVRAHQAN
                                                                                                                                                                                                          EFEKPYFKQLMSFVADERSRHTVYPPADQVYSWTEMCDIQDVKVVILGQDPYHGPNQAHG
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95.2%;
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Pred. No. 1.6e-138;
4; Mismatches 9;
301
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and the
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RESULT 3

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Sequence 5713, Application US/10106698

Publication No. US20030109690A1

GENERAL INFORMATION:
APPLICANT: Ruben et al.

TITLE OF INVENTION: Colon and Colon Cancer Associated Polynucleotides and Poly,
FILE REFERENCE: PA005F1

CURRENT APPLICATION UNMEER: US/10/106,698

CURRENT FILING DATE: 2002-03-27

PRIOR APPLICATION NUMBER: PCT/US00/26524

PRIOR APPLICATION NUMBER: US 60/157,137

PRIOR FILING DATE: 2000-09-28

PRIOR APPLICATION NUMBER: US 60/157,137

PRIOR APPLICATION NUMBER: US 60/163,280

PRIOR FILING DATE: 1999-09-29

PRIOR APPLICATION NUMBER: US 60/163,280

PRIOR FILING DATE: 1999-11-03

NUMBER OF SEQ ID NOS: 8564

SOFTWARE: Patentin Ver. 3.0

SEQ ID NO 5713

LENGTH: 292

TYPE: PRI
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US-10-106-698-5713
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Best Local Similarity
Matches 195; Conservat
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SEQ ID NO 12
LENGTH: 304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 12, Application US/10038010 Publication No. US20030040089A1 GENERAL INFORMATION:
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ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: DNA 91ycosylase
LOCATION: (1)..(304)
OTHER INFORMATION:
ORGANISM: Homo sapiens -10-106-698-5713
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: HYBRIGENICS
APPLICANT: Pierre, Legrain
TITLE OF INVENTION: Protein-protein interactions
FILE REFERENCE: B4767A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      254 DRKRHHVLQAVHÐSPLSAHRGFLGCKHFSKANGLLKLSGTEPINWRAL 301
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        77 AARNVPVGFGESWKKHLSGEFGKPYFIKLMGFVAEERKHYTVYPPPHQVFTWTQMCDIKD
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Query Match

DB 14; Length

292;

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RESULT 6
U8-10-369-493-21107; Sequence 21107, A
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US-10-369-493-236
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, ORGANISM: Xenorhabdus nematophilus
US-10-369-493-236
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PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
SEQ ID NO 236
LENGTH: 225
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
APPLICANTON: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10(52052)B
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
CURRENT FILING DATE: 2003-02-26
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                                                                                                                                      VHPSPLSAHRGFLGCKHFSKANGLLKLSGTEPINW
                                                                                                                                                                                  LTVERGNAHSHANLGWETFTDKVIAAINEHRSGVIFLLWGAHAQKKGRIIDTQRHHVLKA
                                                                                                                                                                                                                   LTVRAHQANSHKDRGWETFTDAVIKWLSVNREGVVFLLWGSYAHKKGATIDRKRHHVLQA 263
                                                                                                                                                                                                                                                             YHGPNQAHGLAFSVQPGIPAPPSLVNMYKELESDIAGFQRPNHGCLLSWXKQGVLLLNTV
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                                                                                                     PHPSPLSAHRGFLGCKHFSQANHLLEAQGLASIDW
  Application US/10369493
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Pred. No. 2.2e-65;
Pred. No. 5.2e-65;
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APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Chen, Xienfeng
ITILE OF INVENTION: EXPRESSION OF MICROBIAL PRO
ITITLE OF INVENTION: PLANTS WITH IMPROVED PROPE
ILLE REFERENCE: 38-10(52052)B
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILLING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
SEQ ID NO 23507
LENGTH: 229
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US-10-369-493-23507
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US-10-369-493-23507
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Best Local Similarity 59.1:
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CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
SEQ ID NO 21107
LENGTH: 225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 23507, Application US/10369493 Publication No. US20030233675A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Publication No. US20030233675A1 GENERAL INFORMATION:
                                                                                 Matches
                                                                                                    Query Match
Best Local
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APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
                                                                                                                                                                                                            TYPE: PRT
                                                                                                    Local Similarity
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                                                                              131; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        66 YHGPNQAHGLAFSVQPGIPAPPSLVNMYKELESDITGFSRPNHGCLVSWAKQGVLLLNTV 125
                                       85
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              YHGPNQAHGLCFSVQKPVPPPPSLVNIYKELCTDIDGFKHPGHGDLSGWAKQGVLLLNAV 203
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LTVERGNAHSHANLGWETFTDKVIAAINEHRHGVIFLLWGSHAQKKGHFINTQQHYVLKA 185
                                   TWRRELAAEFEKPYFKQLMSFYADER-SRHTVYPPADQVYSWTEMCDIQDVKVVILGQDP 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PHPSPLSAHRGFLGCKHFSQANSLLEAQGLTPIDW
TWHDVLAEEKQQPYFLNTLQTVASERQSGVTIYPPQKDVFNAFRFTELGDVKVVILGQDP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          42.2%; Score 678.5; DB 15; Length 59.1%; Pred. No. 8.9e-64; ative 31; Mismatches 56; Indels
                                                                                                  42.2%;
                                                                                 23; Mismatches
                                                                                                    Score 677.5; DB 1
Pred. No. 1.2e-63;
                                                                                                                                                                                                                                                                                                                                                                                                    PROPERTIES
                                                                                                                      DB 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     220
                                                                                   60;
                                                                                   Indels
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                                                                                                                        Length
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                                                                                 1:
                                                                                   Gaps
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RESULT 9
US-10-369-493-524
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US-10-260-877-14
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                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                        Sequence 524, Application US/10369493 Publication No. US20030233675A1
APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION
TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10(52052)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 137
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches 124;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 14, Application US/10260877 Publication No. US20030021813A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Abbott Laboratories
APPLICANT: Chovan, Linda E.
APPLICANT: Hessler, Paul E.
APPLICANT: Hessler, Baul E.
APPLICANT: Hessler, Baul E.
APPLICANT: Reich, Karl A.
APPLICANT: Reich, Karl A.
APPLICANT: REICH, Karl A.
APPLICANT: RESENTIAL BACTERIA GENES AND GENOME
TITLE OF INVENTION: ESSENTIAL GENES'
FILE REFERENCE: 6565.US.Pl
CURRENT FILING DATE: 2002-09-30
CURRENT FILING DATE: 2002-09-30
PRIOR APPLICATION NUMBER: US/09/649,145
PRIOR FILING DATE: 2000-08-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 219
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    204
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       84 ETWRRELAAEFEKPYFKQLMSFVADER-SRHTVYPPADQVYSWTEMCDIQDVKVVILGQD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                APHPSPLSAHRGFFGCRHFSKTNSYLESHGIKPIDWQ 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AVHPSPLSAHRGFLGCKHFSKANGLLKLSGTEPINWR 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PYHGPNQAHGLCFSVQKPVPPPPSLVNIYKELCTDIDGFKHPGHGDLSGWAKQGVLLLNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KNWTDVIGTEKAQPYFQHTLQQVHLARASGKTIYPPQEDVFNAFKYTAFEDVKVVILGQD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 VLTVERGMAHSHANLGWERFTDKVIAVLNEHREKLVFLLWGSHAQKKGQMIDRTRHLVLT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PYHGPNOAHGLAFSVKPEVAIPPSLLNIYKELTQDISGFOMPSNGYLVKWAEQGVLLLNT 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LTVRAGQAHSHASLGWETFTDKVISLINQHREGVVFLLWGSHAQKKGAIIDKQRHHVLKA 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LTVRAHQANSHKDRGWETFTDAVIKWLSVNREGVVFLLWGSYAHKKGATIDRKRHHVLQA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PHPSPLSAHRGFFGCNHFVLANQWLEQRGETPIDW 220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               39.3%; Score 631.5; DB 14; 57.1%; Pred. No. 8.9e-59;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27;
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APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10 (52052) B
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
CURRENT FILING DATE: 2003-02-28
CURRENT FILING DATE: 2003-02-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 15993
LENGTH: 214
                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 15993, Application US/10369493
Publication No. US20030233675A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 124; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                            120
                                                                                                                                   204
                  264 VHPSPLSAHRGFLGCKHFSKANGLLKLSGTEPINW 298
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              139
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                                                                                                       LTVRAHQANSHKDRGWETFTDAVIKWLSVNREGVVFLLWGSYAHKKGATIDRKRHHVLQA
                                                                                                                                                                                                                                                                                                          SWKARVGEWILQPQMQELSAFLRQRKAANARVFPPGPQIFAAFDATPFEQVKVVVLGQDP
                                                                                                                                                                                          YHGEGQAHGLCFSVLPGVPVPPSLLNIYKEIQDDL-GIPRPDHGYLMPWARQGVLLLNAV 119
                                                                                                                                                                                                                                             YHGPNQAHGLCFSVQKPVPPPPPSLVNIYKELCTDIDGFKHPGHGDLSGWAKQGVLLLNAV
                                                                                                                                                                                                                                                                                                                                                                    TWRRELAAEFEKPYFKQLMSFVADERSRHT-VYPPADQVYSWTEMCDIQDVKVVILGQDP 143
                                                                         LTVEQGRAGAHQNKGWEGFTDHVVETLNSEREGLVFLLWGSYAQSKGKVIDQARHRVFKA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HVLQAVHPSPLSAHRGFLGCKHFSKANGLLKLSGTEPINWR 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LLNAVLTVRAHQANSHKDRGWETFTDAVIKWLSVNREGVVFLLWGSYAHKKGATIDRKRH 258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LLNAVLTVRAGQANSHQGKGWEHFTDAVIKAVNAKEERVVFILWGSYARKKKKLITGKNH 199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LGODPYHGPNOAHGLSFSVRPGVRVPPSLRNIYKELTEDIPGFVAPKHGYLRSWAEQGVL 139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LGQDPYHGPNQAHGLCFSVQKPVPPPPSLVNIYKELCTDIDGFKHPGHGDLSGWAKQGVL 198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PANLPEDWQEALLPEFSAPYFHELTDFLRQERKEYTIYPPAPDVFNALRYTPLGEVKVLI
                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       38.4%;
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56.1%; Pred. No. 1.1e-58;
                                                                                                                                                                                                                                                                                                                                                                                                                              35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                           Score 617; DB 15;
Pred. No. 3.1e-57;
5; Mismatches 63;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      70;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 214;
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                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
ITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
ITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10 (52052)B
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT APPLICATION NUMBER: US 60/360,039
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR PILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
SEQ ID NO 15618
LENGTH: 237
TYPE: PRT
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                                          PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
SEQ ID NO 13886
LENGTH: 216
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 13886, Application US/10369493 Publication No. US20030233675A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15618, Application US/10369493 Publication No. US20030233675A1
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Best Local Similarity 53.5%;
  ORGANISM: Pseudomonas fluorescens -10-369-493-13886
                                                                                                                                                                                              APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
TITLE OP INVENTION: EXPRESSION OF MICROBIAL PROTEINS
TITLE OP INVENTION: EXPRESSION OF MICROBIAL PROTEINS
TITLE OP INVENTION: DELOTES WITH IMPROVED PROPERTIES
FILE REPERENCE: 39-10(52052)B
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LTVEQGRAGAHQNKGWEGFTDHVVETLNSEREGLVFLLWGSYAQSKGKVIDQARHRVFKA 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LTVRAHQANSHKDRGWETFTDAVIKWLSVNREGVVFLLWGSYAHKKGAT1DRKRHHVLQA 263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PHPSPLSAHRGFLGCKHFSKTNDHLQRRGLSPIDW
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SWKARVGEWLLQPQMQELSAFLRQRKAANARVFPPGPQIFAAFDATPFEQVKVVVLGODP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 617; DB 15;
Pred. No. 3.5e-57;
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RESULT 14
US-10-369-493-11585
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APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Slater, Steven C.
APPLICANT: Coldman, Barry S.
APPLICANT: Chen, Xianfeng
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES FILE REFERENCE: 38-10(52052)B
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT APPLICATION NUMBER: US-28
PRIOR APPLICATION NUMBER: US-28
PRIOR APPLICATION NUMBER: US-28
PRIOR APPLICATION NUMBER: US-28
PRIOR FILING DATE: 2002-02-21
J. NUMBER OF SEQ ID NOS: 47374
SEQ ID NO 10331
LENGTH: 220
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US-10-369-493-10331
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Best Local S
Matches 114
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                                                                                                                                                                                                                                                                                                                        Matches 112;
                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: unsure
LOCATION: (1)..(220)
OTHER INFORMATION: unsure at all Xaa locations
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Cytophaga hutchinsonii
                                                                                                                                                                                                                                                                                                                                           ocal Similarity
                                                                              123
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  183
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                               HPSPLSAHRGFLGCKHFSKANGLLKLSGTEPINW 298
                                                                              TVEANTPGSHQKKGWETFTDAVIKKVSDEKEHVVFILMGAYAQKKGEIIDREKHLVLKSA 182
                                                                                                             TVRAHQANSHKDRGWETETDAVIKWLSVNREGVVFLLWGSYAHKKGATIDRKRHHVLQAV 264
                                                                                                                                                              HGPGQANGLCFSVNDGIVQPPSLKNIFKEIESDL-GKSAPVSGNLERWANQGVLLLNATL 122
                                                                                                                                                                                                                                          SWKHRIQDEFNKTYFKNLIQFVKEQYKTRVIYPAGKNIFKAFDACPFDAVEVVIIGQDPY
                                                                                                                                                                                                                                                                                  TWRRELAAEFEKPYFKQLMSFVADERSRHTVYPPADQVYSWTEMCDIQDVKVVILGQDPY 144
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HPSPFSAYNGFFGNKHFSKTNAYLKSVGKKEIDW
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Pred. No. 5.1e-57;
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Pred. No. 1.8e-55;
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Sequence 11585, Application US/10369493

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APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10 (52052) B
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
PRIOR FALING DATE: 2003-02-28
PRIOR APPLICATION UMMBER: US 60/360,039
PRIOR APPLICATION OF SEQ ID NOS: 47374
SEQ ID NO 14855
LENGTH: 220
TYPE: PRT
ORGANT
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                                                                     Query Match
Best Local Similarity
Matches 115; Conserv
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LENGTH: 217
                                                                                                                                                         ORGANISM: Agrobacterium tumefaciens -10-369-493-14855
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APPLICANT: Hink
APPLICANT: Slat
APPLICANT: Gold
APPLICANT: Chen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     263 AVHPSPLSAHRGFLGCKHFSKANGLLKLSGTEPINWR 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       180 SPHPSPLSAHNGFFGNGHFSKANAFLVSHGRDPIDWQ 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               120 VLTVBEARAASHQGQGWEKFTDAVIRAVNDECDHVVFLLWGSYAQKKAAFVDQRKHLVLR
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                               84 ETWRRELAAEFEKPYFKQLMSF-VADERSRHTVYPPADQVYSWTEMCDIQDVKVVILGQD 142
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SWKHVLSGEFASPYMQKLKEFLLAEKTAGKRIFPKGAEYFRALDLTFLDEVKVVILGQD 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      : Hinkle, Gregory J.
: Slater, Steven C.
: Goldman, Barry S.
: Chen, Xianfeng
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                                                                               Conservative
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                                                                         37.0%; Score 594; DB 15; 53.0%; Pred. No. 9.1e-55; tive 38; Mismatches 62;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             37.0%;
53.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             38; Mismatches 62;
                                                                                                               Length 220;
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                                                                             Indels
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Search completed: March 18, 2004, 00:11:37 Job time : 37 secs

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Maximum DB seq
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Scoring table:
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                                                                                                                                                                                                                                                                                                                                 score greater than or equal to the score of the result beir and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                 Pred. No.
                1036
693.5
678.5
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617
617
615
600.5
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                                                                                                                                                                                        1550
1421
1036
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1: /cgn2 6/ptcdata/1/pubpaa/USO7_PUBCOMB.pep:*

2: /cgn2 6/ptcdata/1/pubpaa/USO6_NEW_PUB.pep:*

3: /cgn2 6/ptcdata/1/pubpaa/USO6_NEW_PUB.pep:*

4: /cgn2 6/ptcdata/1/pubpaa/USO6_PUBCOMB.pep:*

5: /cgn2 6/ptcdata/1/pubpaa/USO6_PUBCOMB.pep:*

6: /cgn2 6/ptcdata/1/pubpaa/USO6_NEW_PUB.pep:*

7: /cgn2 6/ptcdata/1/pubpaa/USO8_NEW_PUB.pep:*

8: /cgn2 6/ptcdata/1/pubpaa/USO8_NEW_PUB.pep:*

8: /cgn2 6/ptcdata/1/pubpaa/USO8_NEW_PUB.pep:*
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Gapop 10.0 , Gapext 0.5
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(without alignments)
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1 MIGQQHINSFFSPVSKKRVS.....SKANGLLKLSGTEPINWRAL 301
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Maximum Match 100%
Listing first 45 s
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/cgn2_6/ptodata/1/pubpaa/US09C_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US09C_PUBCOMB.pep:*
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/cgn2_6/ptodata/1/pubpaa/US10C_PUBCOMB.pep:*
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 US-10-369-493-236

US-10-369-493-23107

US-10-369-493-23507

US-10-260-877-14

US-10-369-493-1593

US-10-369-493-1593

US-10-369-493-13886

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US-09-758-017A-2
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Million cell updates/sec
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         Sequence 4, Appli
Sequence 2, Appli
Sequence 5713, Appl
Sequence 12, Appl
Sequence 236, Appl
Sequence 21107, A
Sequence 23507, A
Sequence 14, Appl
Sequence 524, Appl
Sequence 1593, A
Sequence 15818, A
Sequence 13886, A
Sequence 11885, A
                                                                                                                                                                                                                                                     Description
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4 5	44	43	42	41	40	9	38	37	36	35	34	3	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16
164	254.5	275	345	359	363.5	380	450.5	459	488.5	493	502.5	502.5	503.5	514		535.5	556.5	558	566	566.5	568.5	571	573.5	575	584.5	587	588	588.5	594
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	24-599-	09-925-30	-10-156-7	-10-369-49	8-62	-09-882-2	-10-369-49	-10-369-493-133	-10-369-493-	-10-424-599	US-10-369-493-9579	-10-369-493	-10-369-493	-10-369-493	-10	-10-289-762-82	-10-369-493-22	-10-369-493-18	-10-156-761-1453	9-493-	69-493-17	-10-369-493-1783	-10-369-493-94	-10-369-493-15	-10-369-493-	US-10-369-493-17754	-10-369-493-91	9-493-166	69-493-14
Ф	ø	1562,	e 9066,	e 8212,	4961,	e 322,	Sequence 18312, A	e 13321	e 3953,	e 1889:	O	e 8693,	e 1912:	e 4570,	e 7327	e 823,	e 2266:	0	e 1453	2335	e 1746!	1783	9434	e 1504	e 1073	e 1775	e 9179,	e 1661	189

ALIGNMENTS

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FILE REFERENCE: U013209-3
CURRENT APPLICATION NUMBER: US/09/758,017A
CURRENT FILING DATE: 2001-01-10
PRIOR APPLICATION NUMBER: 2000 5428
PRIOR FILING DATE: 2000-10-27
PRIOR APPLICATION NUMBER: 2000 0163
PRIOR APPLICATION NUMBER: 2000 0163
PRIOR FILING DATE: 2000-01-12
NUMBER OF SEQ ID NOS: 19
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 4
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                                                                                                                                                                                                                                                                ; LENGTH: 301
; TYPE: PRT
; ORGANISM: Gadus morhua
US-09-758-017A-4
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US-09-758-017A-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION
                                                                                                                                                                       Matches 290;
                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Lanes, Olav
APPLICANT: Willasen, Nils Peder
APPLICANT: Willasen, Nils Peder
APPLICANT: Guddal, Per Henrik
APPLICANT: Guddal, Per Henrik
TITLE OF INVENTION: Cod uracil-DNA glycosylase, gene coding therefore,
TITLE OF INVENTION: recombinant DNA containing said gene or operative parts
TITLE OF INVENTION: thereof, a method for preparing said protein and the
TITLE OF INVENTION: use of said protein or said operative pa
61 RMAKNKKAALDKIRAKATPAGFGETWRRELAAEFEKPYFKQLMSFVADERSRHTVYPPAD
                                                                                              1 MIGQQHINSFFSPVSKKRVSKELGKTEKHAEEVQITPKKLRSSNVEQKTSSPQLSVEQLE
                                                                             MIGQQHINSFFSPVSKKRVSKELGKTEKHAEEVQITPKKLRSSNVEQKTSSPQLSVEQLE
                                                                                                                                                                       Conservative
                                                                                                                                                                                           96.6%;
                                                                                                                                                                                           Score 1550; DB 9;
Pred. No. 4.5e-147;
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                                                                                                                                                                         9
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Sequence

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US-09-758-017A-2
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SEQ ID NO 2
LENGTH: 301
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Lanes, Olav
APPLICANT: Willasen, Nils Peder
APPLICANT: Willasen, Nils Peder
APPLICANT: Guddal, Per Henrik
APPLICANT: Gellesvik, Dag Rune
TITLE OF INVENTION: Cod uracil-DNA glycosylase, gene coding therefore,
TITLE OF INVENTION: recombinant DNA containing said gene or operative
TITLE OF INVENTION: thereof, a method for preparing said protein and t
TITLE OF INVENTION: use of said protein or said operative pa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 2000 0163
PRIOR FILING DATE: 2000-01-12
NUMBER OF SEQ ID NOS: 19
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CURRENT APPLICATION NUMBER: US/09/758,017A
CURRENT FILING DATE: 2001-01-10
                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Gadus morhua
 273
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RGFLGCKHFSKANGLLKLSGTEPINWRAL
                            RGFLGCKHFSKANGLLKLSGTEPINWRAL
                                                                 SHKDRGWETFTDAVIKWLSVNREGVVFLLWGSYAHKKGATIDRKRHHVLQAVHPSPLSAH
                                                                                     SHKDRGWETFTDAVIKWLSVNREGVVFLLWGSYAHKKGATIDRKRHHVLQAVHPSPLSAH
                                                                                                                                   LCFSVQKPVPPPPSLVNIYKELCTDIDGFKHPGHGDLSGWAKQGVLLLNAVLTVRAHQAN
                                                                                                                                                         LCESVQKPVPPPPSLVNIYKELCTDIDGFKHPGHGDLSGWAKQGVLLLNAVLTVRAHQAN
                                                                                                                                                                                                                                     EFEKPYFKQLMSFVADERSRHTVYPPADQVYSWTEMCDIQDVKVVILGQDPYHGPNQAHG
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                                                                                                                                                                                                       EFEKPYFKQLMSFVADERSRHTVYPPADQVYSSTEMCDIQDVKVVILGQDPYHGPNQAHG
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                                                                                                                                                                                                                                                                                                                                                       Score 1421; DB 9;
Pred. No. 4.2e-134;
301
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RESULT 3

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US-10-038-010-12
Sequence 12, Application US/10038010
Publication No. US20030040089A1
GENERAL INFORMATION:
US-10-038-010-12
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                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 60/259,377
PRIOR FILING DATE: 2001-01-02
NUMBER OF SEQ ID NOS: 67
SOFTWARE: Patentin version 3.1
SEQ ID NO 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Ruben et al.
TITLE OF INVENTION: Colon and Colon Cancer Associated Polynucleotides and Polypeptide
FILE REFERENCE: PA005p1
CURRENT APPLICATION NUMBER: US/10/106,698
CURRENT FILING DATE: 2002-03-27
PRIOR APPLICATION NUMBER: PCT/US00/26524
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US 60/157,137
PRIOR APPLICATION NUMBER: US 60/157,137
PRIOR APPLICATION NUMBER: US 60/157,280
PRIOR APPLICATION NUMBER: US 60/163,280
PRIOR FILING DATE: 1999-11-03
NUMBER OF SEQ ID NOS: 8564
SOCTWARE: Patentin Ver. 3.0
SEQ ID NO 5713
LENGTH: 292
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/038,010 CURRENT FILING DATE: 2002-07-23
                                                                                                                                                                                                                                                                                                                       APPLICANT: HYBRIGENICS
APPLICANT: Pierre, Legrain
TITLE OF INVENTION: Protein-protein interactions
FILE REFERENCE: B4767A
                 NAME/KEY: DNA glycosylase LOCATION: (1)..(304) OTHER INFORMATION:
                                                                                LENGTH: 304
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
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ORGANISM: Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LSVYRGFFGCRHFSKTNELLQKSGKKPIDWKEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HQANSHKERGWEQFTDAVVSWLNQNSNGLVFLLWGSYAQKKGSAIDRKRHHVLQTAHPSP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HQANSHKDRGWETFTDAVIKWLSVNREGVVFLLWGSYAHKKGATIDRKRHHVLQAVHPSP 268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   QAHGLCFSVQRPVPPPPSLENIYKELSTDIEDFVHPGHGDLSGWAKQGVLLLNAVLTVRA
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Pred. No. 2e-9
35; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6; DB 14;
2e-95;
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                                                                                                                                                                                                                                                                                                                                             in adipocyte cells
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GENERAL INFORMATION:

APPLICANT: Cao, Yongwei

APPLICANT: Hinkle, Gregory J.

APPLICANT: Slater, Seeven C.

APPLICANT: Slater, Seeven C.

APPLICANT: Goldman, Barry S.

APPLICANT: Goldman, Barry S.

APPLICANT: Chen, Xianfeng

TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF

TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES

FILE REFERENCE: 38-10(52052)B

CURRENT APPLICATION NUMBER: US/10/369,493

CURRENT APPLICATION NUMBER: US/0369,493

CURRENT APPLICATION NUMBER: US 60/360,039

PRIOR PILING DATE: 2002-02-21

NUMBER OF SEQ ID NOS: 47374
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RESULT 6
US-10-369-493-21107
; Sequence 21107, Application US/10369493
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US-10-369-493-236
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US-10-369-493-236
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 43.2%; Score 693.5; DB 15; Length Best Local Similarity 60.5%; Pred. No. 3.7e-61; Matches 130; Conservative 29; Mismatches 55; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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                                                                                                                                                                                               126
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                                                                                                             186
                                                                                                                                                                                                                      204 LTVRAHQANSHKDRGWETFTDAVIKWLSVNREGVVFLLWGSYAHKKGATIDRKRHHVLQA 263
                                                                                                                                                   264 VHPSPLSAHRGFLGCKHFSKANGLLKLSGTEPINW 298
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                                                                                                                                                                                                                                                                                  66
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                                                                                                           PHPSPLSAHRGFLGCKHFSQANHLLEAQGLASIDW
                                                                                                                                                                                             TVERGNAHSHANLGWETFTDKVIAAINEHRSGVIFLLWGAHAQKKGRIIDTQRHHVLKA
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; Pred. No. 2.1e-95;
35; Mismatches 50;
                                                                                                             220
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APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Slater, Steven C.
APPLICANT: Coldman, Barry S.
APPLICANT: Chen, Xianfeng
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES FILE REFERENCE: 38-10(52652)B
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
SEQ ID NO 21107
LENGTH: 225
TYPE: PRT
COCCANTEM: Xenorhabdus nematophilus
                                                                                                                                                                                                                                                       APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Greyen C.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
ITITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN
ITITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10(52052)B
CUURRENT APPLICATION NUMBER: US/10/369,493
CUURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR APPLICATION PRIOR FILING DATE: 2002-02-21
NUMBER OF SEG ID NOS: 47374
SEQ ID NO 23507
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US-10-369-493-23507
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                                                                                                                                                                                           ; TYPE: PRT; ORGANISM: Escherichia coli US-10-369-493-23507
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 23507, Application US/10369493 Publication No. US20030233675A1
                                                                                                                     Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION
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                                                                                                 Matches
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                                                                                               42.2%; Score 677.5; DB 1
Local Similarity 60.9%; Pred. No. 1.5e-59;
hes 131; Conservative 23; Mismatches 60
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1es 127; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          144 YHGBNQAHGLCFSVQKPVPPPPSLVNIYKELCTDIDGFKHPGHGDLSGWAKQGVLLLNAV 203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 66 YHGPNQAHGLAFSVQPGIPAPPSLVNMYKELESDITGFSRPNHGCLVSWAKQGVLLLNTV 125
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LTVRAHQANSHKDRGWETFTDAVIKWLSVNREGVVFLLWGSYAHKKGATIDRKRHHVLQA 263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TWHEVIGHEKTOPYFVDTLTYVANERKAGKTIYPLQQDVFNAFRYTELSDVRVVILGQDP 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TWRRELAAEFEKPYFKQLMSFVADER-SRHTVYPPADQVYSWTEMCDIQDVKVVILGQDP 143
TWHDVLAEEKQQPYFLNTLQTVASERQSGVTIXPPQKDVFNAFRFTELGDVKVVILGQDP
                                          TWRRELAAEFEKPYFKQLMSFVADER-SRHTVYPPADQVYSWTEMCDIQDVKVVILGQDP 143
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                                                                                                                                              DB 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           220
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                                                                                                                                              Length 229;
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US-10-369-493-524
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APPLICANT: Cao, Yongwei
APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianteng
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION (
TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
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US-10-260-877-14
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                                                                                                                                                                                                                                 Sequence 524, Application US/10369493 Publication No. US20030233675A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/260,877
CURRENT FILING DATE: 2002-09-30
PRIOR APPLICATION NUMBER: US/09/649,145
PRIOR FILING DATE: 2000-08-25
NUMBER OF SEQ ID NOS: 137
SOFTWARE: FASESEQ for Windows Version 4.0
SEQ ID NO 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 14, Application US/10260877 Publication No. US20030021813A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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-10-260-877-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Abbott Laboratories
APPLICANT: Chovan, Linda E.
APPLICANT: Hessler, Paul E.
APPLICANT: Hessler, Paul E.
APPLICANT: Reich, Karl A.
TITLE OF INVENTION: ESSENTIAL BACTERIA GENES AND GENOME
TITLE OF INVENTION: SCANNING IN HAEMOPHILUS INFLUENZAE FOR THE IDENTIFICATION OF THE REFERENCE: 6565.US.P1
TILE REFERENCE: 6565.US.P1
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TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                              263
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        62 PYHGPNQAHGLAFSVKPEVAIPPSLLNIYKELTQDISGFQMPSNGYLVKWAEQGVLLLNT 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       84 ETWRRELAABFEKPYFKQLMSFVADER-SRHTVYPPADQVYSWTEMCDIQDVKVVILGQD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 KNWTDVIGTEKAQPYFQHTLQQVHLARASGKTIYPPQEDVFNAFKYTAFEDVKVVILGOD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PYHGPNQAHGLCFSVQKPVPPPPSLVNIYKELCTDIDGFKHPGHGDLSGWAKQGVLLLNA 202
                                                                                                                                                                                                                                                                                                                                                                     APHPSPLSAHRGFFGCRHFSKTNSYLESHGIKPIDWQ 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                VLTVERGMAHSHANLGWERFTDKVIAVLNEHREKLVFLLWGSHAQKKGQMIDRTRHLVLT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              YHGPNQAHGLCFSVQKPVPPPPSLVNIYKELCTDIDGFKHPGHGDLSGWAKQGVLLLNAV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LTVRAGQAHSHASLGWETFTDKVISLINQHREGVVFLLWGSHAQKKGAIIDKQRHHVLKA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27; Mismatches
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GENERAL INCOMPTION:

GENERAL INCOMPTION:

APPLICANT: Cao, Yongwei
APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.

APPLICANT: Slater, Steven C.

APPLICANT: Slater, Steven C.

APPLICANT: Goldman, Barry S.

APPLICANT: Chen, Xianfeng
ITITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF

ITITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES

FILE REFERENCE: 38-10(5252)B

CURRENT FILING DATE: 2003-02-28

CURRENT FILING DATE: 2003-02-28

PRIOR APPLICATION NUMBER: US 60/360,039

PRIOR FILING DATE: 2002-02-21

NUMBER OF SEQ ID NOS: 47374

LENGTH: 214

TYDE: Dem
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; Sequence 15993, Application US/10369493
; Publication No. US20030233675A1
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                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 524
LENGTH: 247
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
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                                              120
264 VHPSPLSAHRGFLGCKHFSKANGLLKLSGTEPINW 298
                                                                                    204 LTVRAHQANSHKDRGWETFTDAVIKWLSVNREGVVFLLWGSYAHKKGATIDRKRHHVLQA
                                                                                                                                                                      144 YHGPNQAHGLCFSVQKPVPPPPSLVNIYKELCTDIDGFKHPGHGDLSGWAKQGVLLLNAV
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                                                                                                                                                                                                                                                                 85 TWRRELAAEFEKPYFKQLMSFVADERSRHT-VYPPADQVYSWTEMCDIQDVKVVILGQDP 143
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                                                                                                                                                                                                                       1 SWKARVGEWLLQPQMQELSAFLRQRKAANARVFPPGPQIFAAFDATPFEQVKVVVLGQDP
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                                          LTVEQGRAGAHQNKGWEGFTDHVVETLNSEREGLVFLLWGSYAQSKGKVIDQARHRVFKA
                                                                                                                                YHGEGQAHGLCFSVLPGVPVPPSLLNIYKEIQDDL-GIPRPDHGYLMPWARQGVLLLNAV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LGQDPYHGPNQAHGLSFSVRPGVRVPPSLRNIYKELTEDIPGFVAPKHGYLRSWAEQGVL
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                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                           38.4%; Score 617; DB 15; 53.5%; Pred. No. 1.7e-53; tive 35; Mismatches 63
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56.1%;
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Pred. No. 7.2e-55;
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APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES FILE REFERENCE: 38-10(52052)B
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
CURRENT FILING DATE: 2003-02-28
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ORGANISM: Pseudomonas fluorescens
US-10-369-493-13886
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES FILE REFERENCE: 38-10(52052)B
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
SEQ ID NO 15618
LENGTH: 237
TYPE: PRI
CONNECTED TO THE PRIOR PR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 13886, Application US/10369493 Publication No. US20030233675A1 GENERAL INFORMATION:
                                                                                                                 PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
SEQ ID NO 13886
LENGTH: 216
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Best Local Similarity 53.5
Matches 115; Conservative
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APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
                                                                                      TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       YHGEGQAHGLCFSVLPGVPVPPSLLNIYKEIQDDL-GIPRPDHGYLMPWARQGVLLLNAV 127
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; Pred. No. 1.9e-53;
35; Mismatches 63;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      222
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 47374
SEQ ID NO 10331
LENGTH: 220
                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
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APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS
TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10 (52052)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: unsure
LOCATION: (1)..(220)
OTHER INFORMATION: unsure at all Xaa locations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Cytophaga hutchinsonii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                    Local Similarity
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183
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                                                                                                                                                                                                                                                                                                                    112;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   144 YHGPNQAHGLCFSVQKDVPPPPSLVNIYKELCTDIDGFKHPGHGDLSGWAKQGVLLLNAV 203
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                                                                                                                                                                                   HGPNQAHGLCFSVQKPVPPPPSLVNIYKELCTDIDGFKHPGHGDLSGWAKQGVLLLNAVL 204
                      HPSPLSAHRGFLGCKHFSKANGLLKLSGTEPINW 298
                                                                                                   TVRAHQANSHKDRGWETFTDAVIKWLSVNREGVVFLLWGSYAHKKGATIDRKRHHVLQAV 264
                                                                                                                                                         HGPGQANGLCFSVNDGIVQPPSLKNIFKEIESDL-GKSAPVSGNLERWANQGVLLLNATL 122
                                                                                                                                                                                                                                       SWKHRIQDEFNKTYFKNLIQFVKEQYKTRVIYPAGKNIFKAFDACPFDAVEVVIIGQDPY 63
                                                                                                                                                                                                                                                                          TWRRELAAEFEKPYFKQLMSFVADERSRHTVYPPADQVYSWTEMCDIQDVKVVILGQDPY 144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MTVERANANAHKDKGWQFFTDRIIEVVSQKQPHLVFMLWGAHAQSKQKLIDATKHLVLTS 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              YHGPGQAHGLCFSVQPGVPAPPSLVNIYKELKRDLN-IDIPNHGYLQSWADQGVLMLNTT 119
HPSPFSAYNGFFGNKHFSKTNAYLKSVGKKEIDW
                                                                            TVEANTPGSHQKKGWETFTDAVIKKVSDEKEHVVFILWGAYAQKKGEIIDREKHLVLKSA 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VHPSPLSAYRGFLGCGHFSRTNKFLEQNGEAPIEWR
                                                                                                                                                                                                                                                                                                                    Conservative
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Pred. No. 2.7e-53;
14; Mismatches 66;
                                                                                                                                                                                                                                                                                                                    Score 600.5; DB 15;
Pred. No. 7.9e-52;
By Mismatches 68;
                                                                                                                                                                                                                                                                                                                                                       DB 15; Length 220;
216
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RESULT 14 US-10-369-493-11585

Sequence 11585, Application US/10369493

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; ORGANISM: Agrobacterium tumefaciens US-10-369-493-14855
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US-10-369-493-14855
Sequence 14855, Application US/10369493
Publication No. US20030233675A1
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; ORGANISM: Agrobacterium tumefaciens
US-10-369-493-11585
                                                                                                                                                                    TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10(52052)8
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
NUMBER: OF SEQ ID NOS: 47374
SEQ ID NO 14855
LENGTH: 220
TYPE: PRT
                                                                    Query Match
Best Local Similarity
                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
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FILE REFERENCE: 38-10(5202)B
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
SEQ ID NO 11585
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APPLICANT: Hinkle, G
APPLICANT: Slater, S
APPLICANT: Goldman,
APPLICANT: Chen, Xić
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Best Local Similarity
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                                                115;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     263 AVHPSPLSAHRGFLGCKHFSKANGLLKLSGTEPINWR 299
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84 ETWRRELAAEFEKPYFKQLMSF-VADERSRHTVYPPADQVYSWTEMCDIQDVKVVILGQD 142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     203 VLTVRAHQANSHKDRGWETFTDAVIKWLSVNREGVVFLLWGSYAHKKGATIDRKRHHVLQ 262
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              143 PYHGPNQAHGLCFSVQKFVPPPPSLVNIYKELCTDIDGFKHPGHGDLSGWAKQGVLLLNA 202
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PYHGLGQAHGLCFSVQPGVRIPPSLVNIYKELQSDL-GIRPVKHGFLESWAKQGVLLLINS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Slater, Steven C. Goldman, Barry S. Chen, Xianfeng
                                          37.0%; Score 594; DB 15; llarity 53.0%; Pred. No. 3.6e-51; Conservative 38; Mismatches 62;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yongwei
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   37.0%; Score 594; DB 15; 53.0%; Pred. No. 3.5e-51; tive 38; Mismatches 62;
                                                                                    Length 220;
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                                              Indels
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